



# SEQUENCE LISTING

<110> Cambridge University Technical Services

<120> A novel family of beta sub-unit proteins from a voltage gated sodium channel nucleic acids encoding them and therapeutic or diagnostic uses thereof

<130> 674558-2001

<140> 09/997,579

<141> 2001-10-15

<150> PCT/EP00/01783

<151> 2000-02-24

<150> 60,129,473

<151> 2000-02-24

<160> 49

<170> PatentIn version 3.2

<210> 1

<211> 215

<212> PRT

<213> Rat

<400> 1

Met Pro Ala Phe Asn Arg Leu Leu Pro Leu Ala Ser Leu Val Leu Ile  
1 5 10 15

Tyr Trp Val Arg Val Cys Phe Pro Val Cys Val Glu Val Pro Ser Glu  
20 25 30

Thr Glu Ala Val Gln Gly Asn Pro Met Lys Leu Arg Cys Ile Ser Cys  
35 40 45

Met Lys Arg Glu Glu Val Glu Ala Thr Thr Val Val Glu Trp Phe Tyr  
50 55 60

Arg Pro Glu Gly Gly Lys Asp Phe Leu Ile Tyr Glu Tyr Arg Asn Gly  
65 70 75 80

His Gln Glu Val Glu Ser Pro Phe Gln Gly Arg Leu Gln Trp Asn Gly  
85 90 95

Ser Lys Asp Leu Gln Asp Val Ser Ile Thr Val Leu Asn Val Thr Leu  
100 105 110

Asn Asp Ser Gly Leu Tyr Thr Cys Asn Val Ser Arg Glu Phe Glu Phe  
115 120 125

Glu Ala His Arg Pro Phe Val Lys Thr Thr Arg Leu Ile Pro Leu Arg  
130 135 140

Val	Thr	Glu	Glu	Ala	Gly	Glu	Asp	Phe	Thr	Ser	Val	Val	Ser	Glu	Ile
145					150					155					160
Met	Met	Tyr	Ile	Leu	Leu	Val	Phe	Leu	Thr	Leu	Trp	Leu	Phe	Ile	Glu
				165					170					175	
Met	Ile	Tyr	Cys	Tyr	Arg	Lys	Val	Ser	Lys	Ala	Glu	Glu	Ala	Ala	Gln
			180					185					190		
Glu	Asn	Ala	Ser	Asp	Tyr	Leu	Ala	Ile	Pro	Ser	Glu	Asn	Lys	Glu	Asn
	195						200					205			
Ser	Val	Val	Pro	Val	Glu	Glu									
	210					215									

<210> 2  
 <211> 215  
 <212> PRT  
 <213> Homo sapiens

<400> 2

Met	Pro	Ala	Phe	Asn	Arg	Leu	Phe	Pro	Leu	Ala	Ser	Leu	Val	Leu	Ile
1				5					10					15	
Tyr	Trp	Val	Ser	Val	Cys	Phe	Pro	Val	Cys	Val	Glu	Val	Pro	Ser	Glu
			20					25					30		
Thr	Glu	Ala	Val	Gln	Gly	Asn	Pro	Met	Lys	Leu	Arg	Cys	Ile	Ser	Cys
		35					40					45			
Met	Lys	Arg	Glu	Glu	Val	Glu	Ala	Thr	Thr	Val	Val	Glu	Trp	Phe	Tyr
	50					55					60				
Arg	Pro	Glu	Gly	Gly	Lys	Asp	Phe	Leu	Ile	Tyr	Glu	Tyr	Arg	Asn	Gly
65					70					75					80
His	Gln	Glu	Val	Glu	Ser	Pro	Phe	Gln	Gly	Arg	Leu	Gln	Trp	Asn	Gly
				85					90					95	
Ser	Lys	Asp	Leu	Gln	Asp	Val	Ser	Ile	Thr	Val	Leu	Asn	Val	Thr	Leu
			100					105					110		
Asn	Asp	Ser	Gly	Leu	Tyr	Thr	Cys	Asn	Val	Ser	Arg	Glu	Phe	Glu	Phe
		115					120					125			
Glu	Ala	His	Arg	Pro	Phe	Val	Lys	Thr	Thr	Arg	Leu	Ile	Pro	Leu	Arg
	130					135					140				
Val	Thr	Glu	Glu	Ala	Gly	Glu	Asp	Phe	Thr	Ser	Val	Val	Ser	Glu	Ile
145					150					155					160
Met	Met	Tyr	Ile	Leu	Leu	Val	Phe	Leu	Thr	Leu	Trp	Leu	Leu	Ile	Glu
				165					170					175	
Met	Ile	Tyr	Cys	Tyr	Arg	Lys	Val	Ser	Lys	Ala	Glu	Glu	Ala	Ala	Gln

	180		185		190
Glu Asn Ala Ser Asp Tyr Leu Ala Ile Pro Ser Glu Asn Lys Glu Asn					
195		200		205	
Ser Ala Val Pro Val Glu Glu					
210		215			

<210> 3  
 <211> 2220  
 <212> DNA  
 <213> rat

<400> 3  
 cgtggccctg gagagggacg gttttgacca cctaatacgtc cagcatcggg gcttcgcaag 60  
 atccaggaac gcgccccacg gaaaggggtc cctcgggtcta cccatcctcc acctctgaga 120  
 tccccaccc caccggaggt cccacctctt tccacccttg aaggacctcc tgtgagcccg 180  
 ggaccctgtg tacaggactg aagtggaaca aattctgtag cccagacgac ggctggagtg 240  
 gggacacgcc caactgaaga agcctgcccc gccgtagaag cccgagatcc tgagtctcgg 300  
 tggattgaag tcgttgtccc tgggggaggc aagagcttca gaaatcgctt acggtggaaa 360  
 agatgcctgc cttcaacaga ttgcttcccc tagcttctct agtgctcatc tactgggtca 420  
 gagtctgctt ccctgtgtgt gtggaagtgc cctcggagac agaagcgggtg cagggcaatc 480  
 ccatgaagct gaggtgcatc tcctgcatga agagggagga ggtggaggcc accactgtgg 540  
 tggagtgggt ctacaggcct gagggcggtg aagatttcct tatatatgag tatcggaatg 600  
 gccaccagga agtggagagc cccttccaag gccgtctgca gtggaatggg agcaaagacc 660  
 tgcaggacgt atccatcact gtactcaatg tcactttgaa tgactctggc ctctacacat 720  
 gcaatgtgtc cagggagttc gaattcgagg cacacaggcc ttttgtgaag accacgagac 780  
 tgataccttt gcgagtcact gaagaggcgg gagaagactt cacctccgtg gtctcggaaa 840  
 tcatgatgta catcctcctg gtcttcctca ccttgtggct gtttattgag atgatctatt 900  
 gctacagaaa ggtctctaag gccgaagagg cagcacagga aaatgcgtct gactaccttg 960  
 ctatcccttc agagaacaag gagaactctg tggtacctgt ggaggaataa tgtggtgtga 1020  
 cttgaggtga tgtacacagg catctgggag ggtgatctga gtgctgaggg actggatatc 1080  
 cccagttcag tgatgccagc aatatcagga agtgccccag gtgtcccaac acatccatct 1140  
 tttctattca tcaaccacca acccaatgtg agattttcac ctgacttccg aactctatca 1200  
 gaactctaca catctttacc ttgcctgaac cgaagagcca acatctatct ctacacggac 1260

taaacctcac tctgttcttg cttccaacca agtaactccc aacttaacta gagttgttcc	1320
ctatgttcca aatgatttag acaagtactg gagagtagta ttacctctgc cctgactgtc	1380
tgtgactggg tcattctcca ctgcagcaaa aggatggata taaatcggaa gaaagccctg	1440
actagtttgt cttaaagcca aagcgtgcc agtacgtact ttgattcatt gaagtcagtt	1500
tttcctgctt ctcagagcgc cagaaagcat gcccctaattg cttgcaggga catcatctgt	1560
gtgcactgga acgctttctg gagctcagtg tttggaggct gtatcccat aatcctgaag	1620
acctggagca aaccagaac ttccaggaag tcccaaggaa ggatccagga cagtttcagg	1680
gtctcgaaaa tgatataaca cactcctgat attggaaaca tggatgagtg acctttctgg	1740
attgaaactc ctcagttctt catgtctcag tgtctgtgga tcagtattag tcctcgtttt	1800
acaggaggaa actgagactc acacaaggct gaacaggaca ttaggggat taaactgggc	1860
cagagatgac tttcctgcc ccaacctcac actccctggg atgagaggta tttttgagga	1920
ctctaacatt cagcatgcc tttgccagc ggaagctgac tgccacagat ctgaggaact	1980
ggaaaccagg taagaaaaca cagacggcat gagatagact tcaggatttc acacaaagat	2040
ttgtgaatct gaagcatcct ccaggagaga cggcacccga gggcaatatc tctgtgatga	2100
aaaatggttt tagtctgaaa tggacagtca acagagagac aaagatgggc gtgtagcttc	2160
taaatacctc acctgtagat gtcacgtttg ggtactggtg tttgtaaagt cccccacctg	2220

<210> 4  
 <211> 1261  
 <212> DNA  
 <213> Homo sapiens

<400> 4	
ccctcccttc cgagctgagc ttaccctggg cgaaacgag cgaggcaggg gcgcgagtgg	60
aagctggagt tccgggggtg gcggggaggc gactgtccgt ggtgctgagc gccggcgaga	120
gcgggcgcgg agcggctgat cggctccctc gaactgggga ggtccagtgg ggtcgcttag	180
ggcccaaagc cccacccgg ctccaaaagc tcccagggcc tcccaggca ccggtgctcg	240
gcccttcctt cggtcagaaa gtcgccccct gggggcagtt cgtcccaaag ggtttcctcg	300
aaagaatctg agagggcgca gtccttgacc gagggaatct ctctgtgtag ccttggaagc	360
cgccagcccc agaagatgcc tgccttcaat agattgtttc ccctggcttc tctcgtgctt	420
atctactggg tcagtgtctg cttccctgtg tgtgtggaag tgccctcgga gacggaggcc	480
gtgcagggca accccatgaa gctgcgctgc atctcctgca tgaagagaga ggaggtggag	540

```

gccaccacgg tgggtggaatg gttctacagg cccgagggcg gtaaagattt ccttatttac 600
gagtatcgga atggccacca ggaggtggag agcccccttc aggggcgcct gcagtggaat 660
ggcagcaagg acctgcagga cgtgtccatc actgtgctca acgtcactct gaacgactct 720
ggcctctaca cctgcaatgt gtcccgggag tttgagtttg aggcgcatcg gccctttgtg 780
aagacgacgc ggctgatccc cctaagagtc accgaggagg ctggagagga cttcacctct 840
gtggtctcag aaatcatgat gtacatcctt ctggtcttcc tcaccctgtg gctgctcatc 900
gagatgatat attgctacag aaaggtctca aaagccgaag aggcagccca agaaaacgcg 960
tctgactacc ttgccatccc atctgagaac aaggagaact ctgcggtacc agtggaggaa 1020
tagaacagga gcagtgtgac atgaggtggc ctgaacacct gagggactgg acatcccatg 1080
ttcagcaatg tcaatggcat caggagggcg cccaagggc cccatcgctt cccttcatgc 1140
atccattggt ctgttcattc attcatccat acatccacct gcctctgagc tttcacctct 1200
gactccctaa ctccatcaga cctctacgca ccataagact ctgccagaac tgagaagccg 1260
g 1261

```

```

<210> 5
<211> 24
<212> PRT
<213> Homo sapiens

```

```
<400> 5
```

```

Met Pro Ala Phe Asn Arg Leu Phe Pro Leu Ala Ser Leu Val Leu Ile
1           5           10           15

```

```

Tyr Trp Val Ser Val Cys Phe Pro
                20

```

```

<210> 6
<211> 24
<212> PRT
<213> rat

```

```
<400> 6
```

```

Met Pro Ala Phe Asn Arg Leu Leu Pro Leu Ala Ser Leu Val Leu Ile
1           5           10           15

```

```

Tyr Trp Val Arg Val Cys Phe Pro
                20

```

```

<210> 7
<211> 19

```

<212> PRT  
<213> homo sapiens

<400> 7

Met Pro Ala Phe Asn Arg Leu Phe Pro Leu Ala Ser Leu Val Leu Ile  
1 5 10 15

Tyr Trp Val

<210> 8  
<211> 19  
<212> PRT  
<213> rat

<400> 8

Met Pro Ala Phe Asn Arg Leu Leu Pro Leu Ala Ser Leu Val Leu Ile  
1 5 10 15

Tyr Trp Val

<210> 9  
<211> 12  
<212> PRT  
<213> homo sapiens

<400> 9

Met Pro Ala Phe Asn Arg Leu Phe Pro Leu Ala Ser  
1 5 10

<210> 10  
<211> 12  
<212> PRT  
<213> rat

<400> 10

Met Pro Ala Phe Asn Arg Leu Leu Pro Leu Ala Ser  
1 5 10

<210> 11  
<211> 15  
<212> PRT  
<213> homo sapiens

<400> 11

Phe Pro Leu Ala Ser Leu Val Leu Ile Tyr Trp Val Ser Val Cys  
1 5 10 15

<210> 12  
<211> 15  
<212> PRT  
<213> rat

<400> 12

Leu Pro Leu Ala Ser Leu Val Leu Ile Tyr Trp Val Arg Val Cys  
1 5 10 15

<210> 13  
<211> 5  
<212> PRT  
<213> homo sapiens

<400> 13

Ser Val Cys Phe Pro  
1 5

<210> 14  
<211> 5  
<212> PRT  
<213> rat

<400> 14

Arg Val Cys Phe Pro  
1 5

<210> 15  
<211> 11  
<212> PRT  
<213> Homo sapiens

<400> 15

Val Leu Ile Tyr Trp Val Ser Val Cys Phe Pro  
1 5 10

<210> 16  
<211> 11  
<212> PRT  
<213> rat

<400> 16

Val Leu Ile Tyr Trp Val Arg Val Cys Phe Pro  
1 5 10

<210> 17  
<211> 39  
<212> PRT  
<213> homo sapiens

<400> 17

Met Pro Ala Phe Asn Arg Leu Phe Pro Leu Ala Ser Leu Val Leu Ile  
1 5 10 15

Tyr Trp Val Ser Val Cys Phe Pro Val Cys Val Glu Val Pro Ser Glu  
20 25 30

Thr Glu Ala Val Gln Gly Asn  
35

<210> 18  
<211> 39  
<212> PRT  
<213> rat

<400> 18

Met Pro Ala Phe Asn Arg Leu Leu Pro Leu Ala Ser Leu Val Leu Ile  
1 5 10 15

Tyr Trp Val Arg Val Cys Phe Pro Val Cys Val Glu Val Pro Ser Glu  
20 25 30

Thr Glu Ala Val Gln Gly Asn  
35

<210> 19  
<211> 9  
<212> PRT  
<213> Homo sapiens

<400> 19

Cys Val Glu Val Pro Ser Glu Thr Glu  
1 5

<210> 20  
<211> 17  
<212> PRT  
<213> Homo sapiens

<400> 20

Trp Val Ser Val Cys Phe Pro Val Cys Val Glu Val Pro Ser Glu Thr  
1 5 10 15

Glu



<210> 21  
<211> 17  
<212> PRT  
<213> Rat

<400> 21

Trp	Val	Arg	Val	Cys	Phe	Pro	Val	Cys	Val	Glu	Val	Pro	Ser	Glu	Thr
1				5					10					15	

Glu

<210> 22  
<211> 159  
<212> PRT  
<213> Homo sapiens

<400> 22

Met	Pro	Ala	Phe	Asn	Arg	Leu	Phe	Pro	Leu	Ala	Ser	Leu	Val	Leu	Ile
1				5					10					15	

Tyr	Trp	Val	Ser	Val	Cys	Phe	Pro	Val	Cys	Val	Glu	Val	Pro	Ser	Glu
		20						25					30		

Thr	Glu	Ala	Val	Gln	Gly	Asn	Pro	Met	Lys	Leu	Arg	Cys	Ile	Ser	Cys
	35						40					45			

Met	Lys	Arg	Glu	Glu	Val	Glu	Ala	Thr	Thr	Val	Val	Glu	Trp	Phe	Tyr
	50						55				60				

Arg	Pro	Glu	Gly	Gly	Lys	Asp	Phe	Leu	Ile	Tyr	Glu	Tyr	Arg	Asn	Gly
65					70					75				80	

His	Gln	Glu	Val	Glu	Ser	Pro	Phe	Gln	Gly	Arg	Leu	Gln	Trp	Asn	Gly
			85						90					95	

Ser	Lys	Asp	Leu	Gln	Asp	Val	Ser	Ile	Thr	Val	Leu	Asn	Val	Thr	Leu
			100					105					110		

Asn	Asp	Ser	Gly	Leu	Tyr	Thr	Cys	Asn	Val	Ser	Arg	Glu	Phe	Glu	Phe
		115					120					125			

Glu	Ala	His	Arg	Pro	Phe	Val	Lys	Thr	Thr	Arg	Leu	Ile	Pro	Leu	Arg
	130					135					140				

Val	Thr	Glu	Glu	Ala	Gly	Glu	Asp	Phe	Thr	Ser	Val	Val	Ser	Glu
145					150					155				

<210> 23

<211> 159  
<212> PRT  
<213> Rat

<400> 23

Met Pro Ala Phe Asn Arg Leu Leu Pro Leu Ala Ser Leu Val Leu Ile  
1 5 10 15  
Tyr Trp Val Arg Val Cys Phe Pro Val Cys Val Glu Val Pro Ser Glu  
20 25 30  
Thr Glu Ala Val Gln Gly Asn Pro Met Lys Leu Arg Cys Ile Ser Cys  
35 40 45  
Met Lys Arg Glu Glu Val Glu Ala Thr Thr Val Val Glu Trp Phe Tyr  
50 55 60  
Arg Pro Glu Gly Gly Lys Asp Phe Leu Ile Tyr Glu Tyr Arg Asn Gly  
65 70 75 80  
His Gln Glu Val Glu Ser Pro Phe Gln Gly Arg Leu Gln Trp Asn Gly  
85 90 95  
Ser Lys Asp Leu Gln Asp Val Ser Ile Thr Val Leu Asn Val Thr Leu  
100 105 110  
Asn Asp Ser Gly Leu Tyr Thr Cys Asn Val Ser Arg Glu Phe Glu Phe  
115 120 125  
Glu Ala His Arg Pro Phe Val Lys Thr Thr Arg Leu Ile Pro Leu Arg  
130 135 140  
Val Thr Glu Glu Ala Gly Glu Asp Phe Thr Ser Val Val Ser Glu  
145 150 155

<210> 24  
<211> 10  
<212> PRT  
<213> Homo sapiens

<400> 24

Thr Thr Arg Leu Ile Pro Leu Arg Val Thr  
1 5 10

<210> 25  
<211> 13  
<212> PRT  
<213> Homo sapiens

<400> 25

Cys Met Lys Arg Glu Glu Val Glu Ala Thr Thr Val Val  
1 5 10

<210> 26  
<211> 10  
<212> PRT  
<213> Homo sapiens

<400> 26

Tyr Glu Tyr Arg Asn Gly His Gln Glu Val  
1 5 10

<210> 27  
<211> 12  
<212> PRT  
<213> Homo sapiens

<400> 27

Trp Asn Gly Ser Lys Asp Leu Gln Asp Val Ser Ile  
1 5 10

<210> 28  
<211> 14  
<212> PRT  
<213> Homo sapiens

<400> 28

Ser Arg Glu Phe Glu Phe Glu Ala His Arg Pro Phe Val Lys  
1 5 10

<210> 29  
<211> 9  
<212> PRT  
<213> Homo sapiens

<400> 29

Val Glu Ser Pro Phe Gln Gly Arg Leu  
1 5

<210> 30  
<211> 13  
<212> PRT  
<213> Homo sapiens

<400> 30

Glu Glu Ala Gly Glu Asp Phe Thr Ser Val Val Ser Glu  
1 5 10

<210> 31  
<211> 34  
<212> PRT  
<213> Homo sapiens

<400> 31

Arg Lys Val Ser Lys Ala Glu Glu Ala Ala Gln Glu Asn Ala Ser Asp  
1 5 10 15

Tyr Leu Ala Ile Pro Ser Glu Asn Lys Glu Asn Ser Ala Val Pro Val  
20 25 30

Glu Glu

<210> 32  
<211> 7  
<212> PRT  
<213> Homo sapiens

<400> 32

Asp Tyr Leu Ala Ile Pro Ser  
1 5

<210> 33  
<211> 22  
<212> DNA  
<213> Artificial sequence

<220>

<223> primer used to amplify nucleic acid sequences encoding b3 subunit  
nucleic acid of rat or human

<400> 33  
atgcctgcct tcaacagatt gc 22

<210> 34  
<211> 20  
<212> DNA  
<213> Artificial sequence

<220>

<223> primer used to amplify nucleic acid sequences encoding b3 subunit  
nucleic acid of rat or human

<400> 34  
ttattcctcc acaggtacca 20

<210> 35  
<211> 45  
<212> DNA

<213> Artificial sequence

<220>

<223> antisense radiolabelled oligonucleotide probe used in the in situ hybridization experiments

<400> 35

ggggaagcaa tctgttgaag gcaggcatct tttccaccgt aagcg

45

<210> 36

<211> 18

<212> DNA

<213> Artificial sequence

<220>

<223> primer used to amplify nucleic acid encoding a b3 sub-unit from a voltage-gated sodium channel

<400> 36

ggtgaagcaa tatggccg

18

<210> 37

<211> 18

<212> DNA

<213> Artificial sequence

<220>

<223> reverse primer (nucleotides 1317-1300) corresponding to unique sequence in the 3' untranslated region of each b subunit

<400> 37

agatgaggcc cagaaccc

18

<210> 38

<211> 20

<212> DNA

<213> Artificial sequence

<220>

<223> forward primer (nucleotides 1942-1961) corresponding to unique sequence in the 3' untranslated region of each b subunit

<400> 38

ggaagctgac tgccacagat

20

<210> 39

<211> 20

<212> DNA

<213> Artificial sequence

<220>

<223> reverse primer (nucleotides 2209-2190) corresponding to unique sequence in the 3' untranslated region of each b subunit

<400> 39  
 cctgggggac tttacaaaca 20

<210> 40  
 <211> 19  
 <212> DNA  
 <213> Artificial sequence

<220>  
 <223> a-tubulin forward primer (nucleotides 298-316) corresponding to  
 unique sequence in the 3' untranslated region of each b subunit

<400> 40  
 cactggtacg tgggtgagg 19

<210> 41  
 <211> 22  
 <212> DNA  
 <213> Artificial sequence

<220>  
 <223> reverse primer (nucleotides 469-448) corresponding to unique  
 sequence in the 3' untranslated region of each b subunit

<400> 41  
 tttgacatga tacagggact gc 22

<210> 42  
 <211> 44  
 <212> DNA  
 <213> Artificial sequence

<220>  
 <223> rat b1 (nucleotides 1296-1252) primer used in immobilization of  
 nucleic acid probe on a substrate

<400> 42  
 gcttgatggg gtgaagaggg gtcgggacag ggacagtagt gggc 44

<210> 43  
 <211> 45  
 <212> DNA  
 <213> Artificial sequence

<220>  
 <223> rat a IIA (nucleotides 1659-1615) primer used in immobilization  
 of nucleic acid probe on a substrate

<400> 43  
 gcagaatcca gagacttcag cggggcaggc gggataggtg ttttc 45

<210> 44  
<211> 218  
<212> PRT  
<213> Rat

<400> 44

Met	Gly	Thr	Leu	Leu	Ala	Leu	Val	Val	Gly	Ala	Val	Leu	Val	Ser	Ser	
1			5						10					15		
Ala	Trp	Gly	Gly	Cys	Val	Glu	Val	Asp	Ser	Glu	Thr	Glu	Ala	Val	Tyr	
		20						25					30			
Gly	Met	Thr	Phe	Lys	Ile	Leu	Cys	Ile	Ser	Cys	Lys	Arg	Arg	Ser	Glu	
		35					40					45				
Thr	Thr	Ala	Glu	Thr	Phe	Thr	Glu	Trp	Thr	Phe	Arg	Gln	Lys	Gly	Thr	
	50					55					60					
Glu	Glu	Phe	Val	Lys	Ile	Leu	Arg	Tyr	Glu	Asn	Glu	Val	Leu	Gln	Leu	
65					70					75					80	
Glu	Glu	Asp	Glu	Arg	Phe	Glu	Gly	Arg	Val	Val	Trp	Asn	Gly	Ser	Arg	
			85						90					95		
Gly	Thr	Lys	Asp	Leu	Gln	Asp	Leu	Ser	Ile	Phe	Ile	Thr	Asn	Val	Thr	
			100					105					110			
Tyr	Asn	His	Ser	Gly	Asp	Tyr	Glu	Cys	His	Val	Tyr	Arg	Leu	Leu	Phe	
		115					120					125				
Phe	Asp	Asn	Tyr	Glu	His	Asn	Thr	Ser	Val	Val	Lys	Lys	Ile	His	Leu	
	130					135					140					
Glu	Val	Val	Asp	Lys	Ala	Asn	Arg	Asp	Met	Ala	Ser	Ile	Val	Ser	Glu	
145					150					155					160	
Ile	Met	Met	Tyr	Val	Leu	Ile	Val	Val	Leu	Thr	Ile	Trp	Leu	Val	Ala	
				165					170					175		
Glu	Met	Val	Tyr	Cys	Tyr	Lys	Lys	Ile	Ala	Ala	Ala	Thr	Glu	Ala	Ala	
			180					185					190			
Ala	Gln	Glu	Asn	Ala	Ser	Glu	Tyr	Leu	Ala	Ile	Thr	Ser	Glu	Ser	Lys	
		195					200					205				
Glu	Asn	Cys	Thr	Gly	Val	Gln	Val	Ala	Glu							
	210					215										

<210> 45  
<211> 119  
<212> PRT  
<213> Rat

<400> 45

Ile Val Val Tyr Thr Asp Arg Glu Val Tyr Gly Ala Val Gly Ser Gln  
 1 5 10 15  
 Val Thr Leu His Cys Ser Phe Trp Ser Ser Glu Trp Val Ser Asp Asp  
 20 25 30  
 Ile Ser Phe Thr Trp Arg Tyr Gln Pro Glu Gly Gly Arg Asp Ala Ile  
 35 40 45  
 Ser Ile Phe His Tyr Ala Lys Gly Gln Pro Tyr Ile Asp Glu Val Gly  
 50 55 60  
 Thr Phe Lys Glu Arg Ile Gln Trp Val Gly Asp Pro Ser Trp Lys Asp  
 65 70 75 80  
 Gly Ser Ile Val Ile His Asn Leu Asp Tyr Ser Asp Asn Gly Thr Phe  
 85 90 95  
 Thr Cys Asp Val Lys Asn Pro Pro Asp Ile Val Gly Lys Thr Ser Gln  
 100 105 110  
 Val Thr Leu Tyr Val Phe Glu  
 115

<210> 46  
 <211> 8  
 <212> PRT  
 <213> Homo sapiens

<400> 46

Glu Gly Gly Lys Asp Phe Leu Ile  
 1 5

<210> 47  
 <211> 34  
 <212> PRT  
 <213> Rat

<400> 47

Arg Lys Val Ser Lys Ala Glu Glu Ala Ala Gln Glu Asn Ala Ser Asp  
 1 5 10 15  
 Tyr Leu Ala Ile Pro Ser Glu Asn Lys Glu Asn Ser Val Val Pro Val  
 20 25 30

Glu Glu

<210> 48  
 <211> 265  
 <212> DNA  
 <213> Rattus norvegicus



```

<400> 48
tcaggaagtg ccccaggtgt cccaacacat ccattctttc tattcatcaa ccaccaaccc 60
aatgtgagat tttcacctga cttccgaact ctatcagaac tctacacatc tttaccttgc 120
ctgaaccgaa gagccaacat ctatctctac acggactaaa cctcactctg ttcttgcttc 180
caaccaagta actcccaact taactagagt tgttccctat gttccaaatg atttagacaa 240
gtactggaga gtagtattac ctctg 265

```

```

<210> 49
<211> 471
<212> DNA
<213> homo sapiens

```

```

<220>
<221> misc_feature
<222> (138)..(138)
<223> n is a, c, g, or t

```

```

<400> 49
gaattcggcc ttcattggcct accagaagat gcctgccttc aatagattgt tccccctggc 60
ttctctcgtg cttatctact gggtcagtgt ctgcttcctt gtgtgtgtgg aagtgccttc 120
ggagacggag ccgtgcangg caaccccatg aagctgcgct gcatctcctg catgaagaga 180
gaggaggtgg aggccaccac ggtggtggaa tggttctaca ggcccagggg cggtaaagat 240
ttccttattt acgagtatcg gaatggccac caggaggtgg agagcccctt tcaggggcgc 300
ctgcagtggg atggcagcaa ggacctgcag gacgtgtcca tcaactgtgct caacgtcact 360
ctgaacgact ctggcctcta cacctgcaat gtgtcccggg agtttgagtt tgaggcgcat 420
cggccctttg tgaagacgac gcggctgac cccctaagag tcggactcga g 471

```